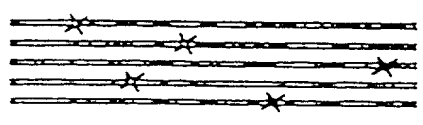
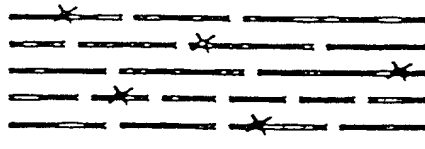


Figure 1 A

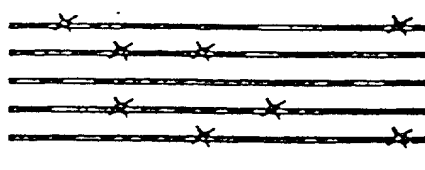
Family of genes {



Fragmentation
DNase I {



PCR
without primers {



PCR
with primers {




Figure 1 B

(Double-stranded process carried out, but illustrated here with a single strand)

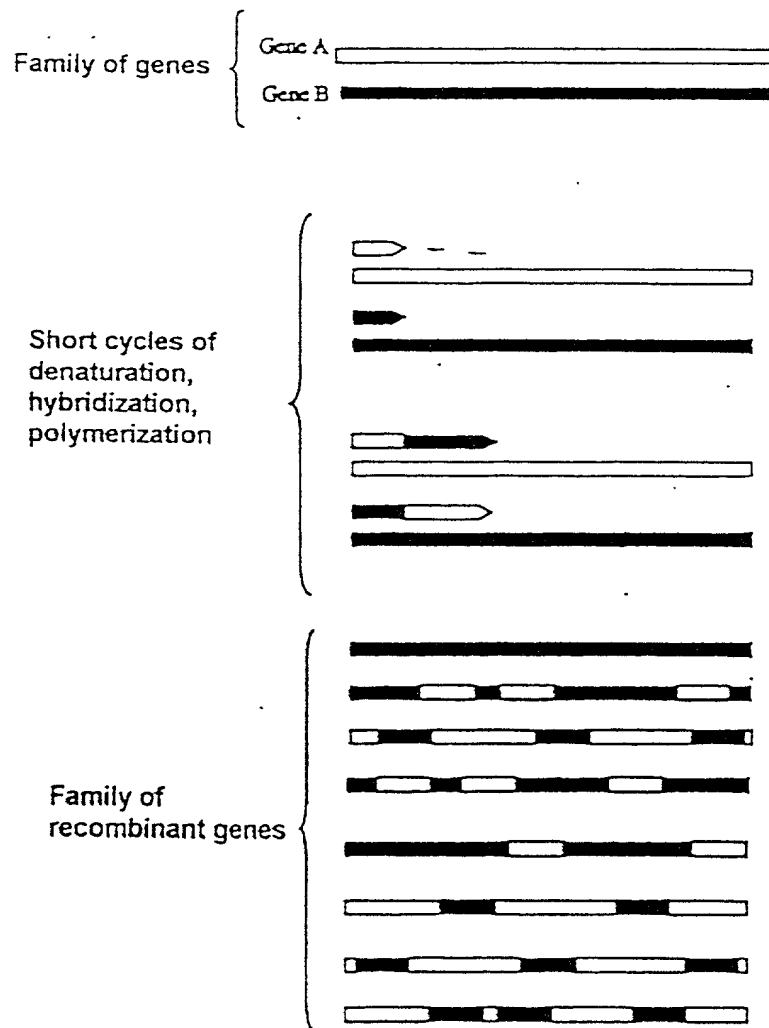


Fig.2

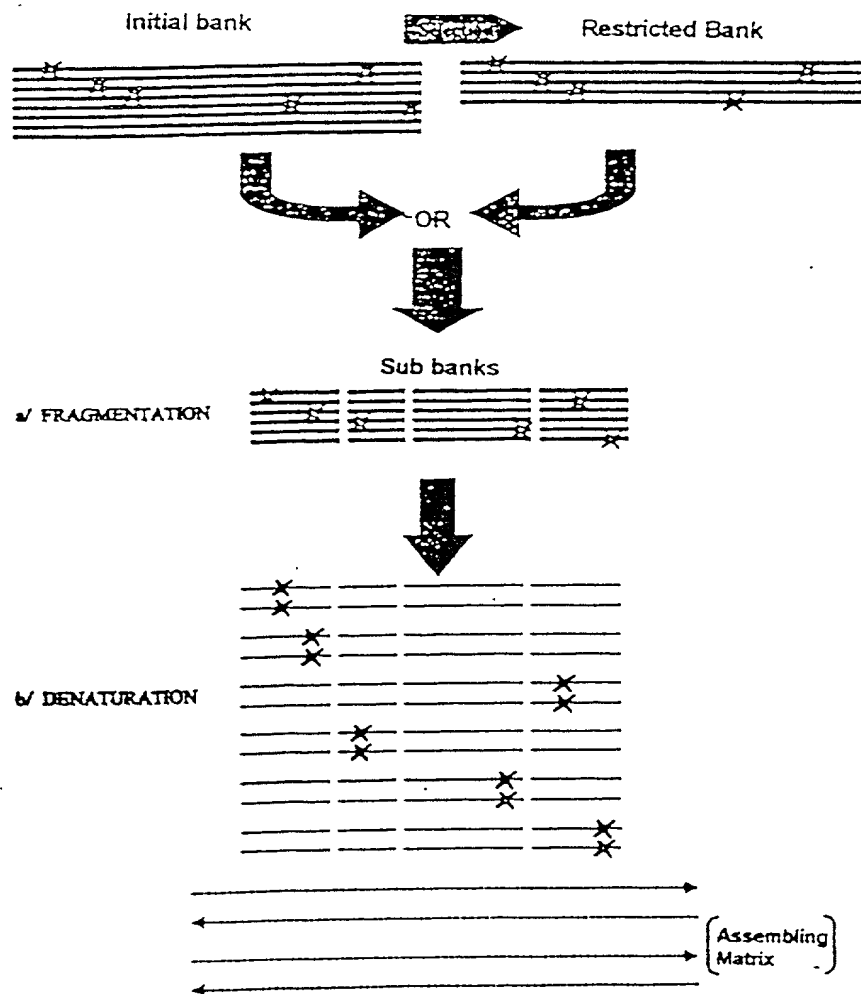


Fig. 2- continued

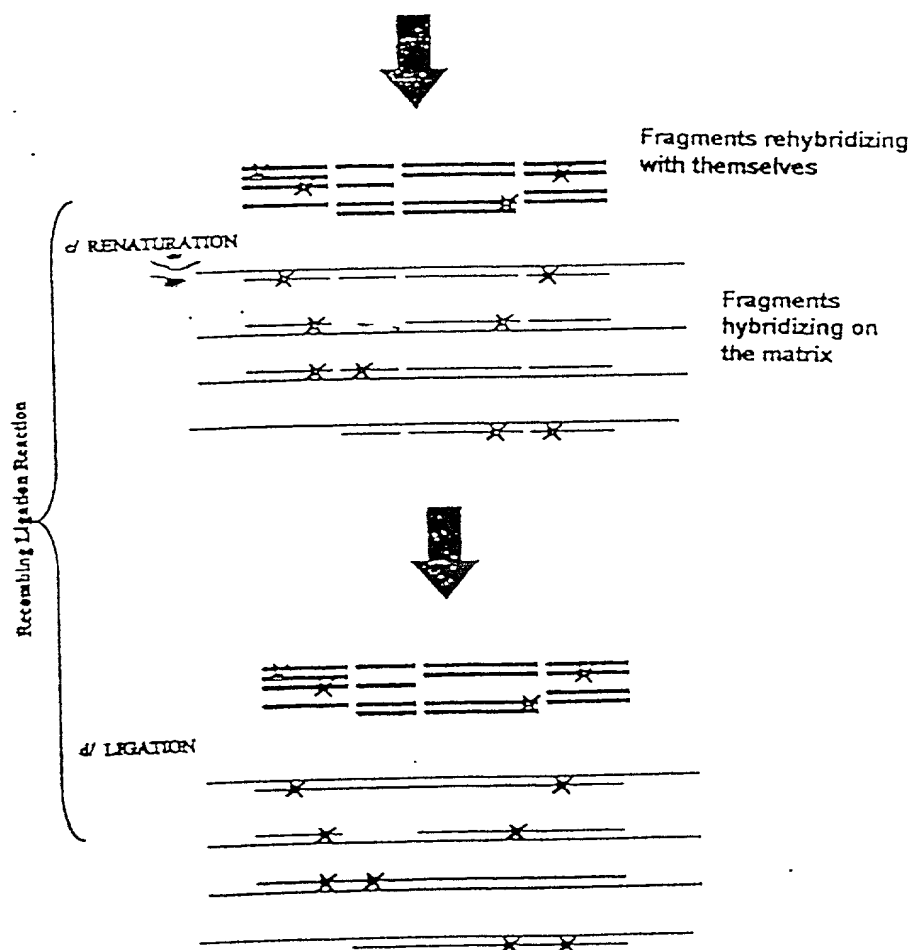
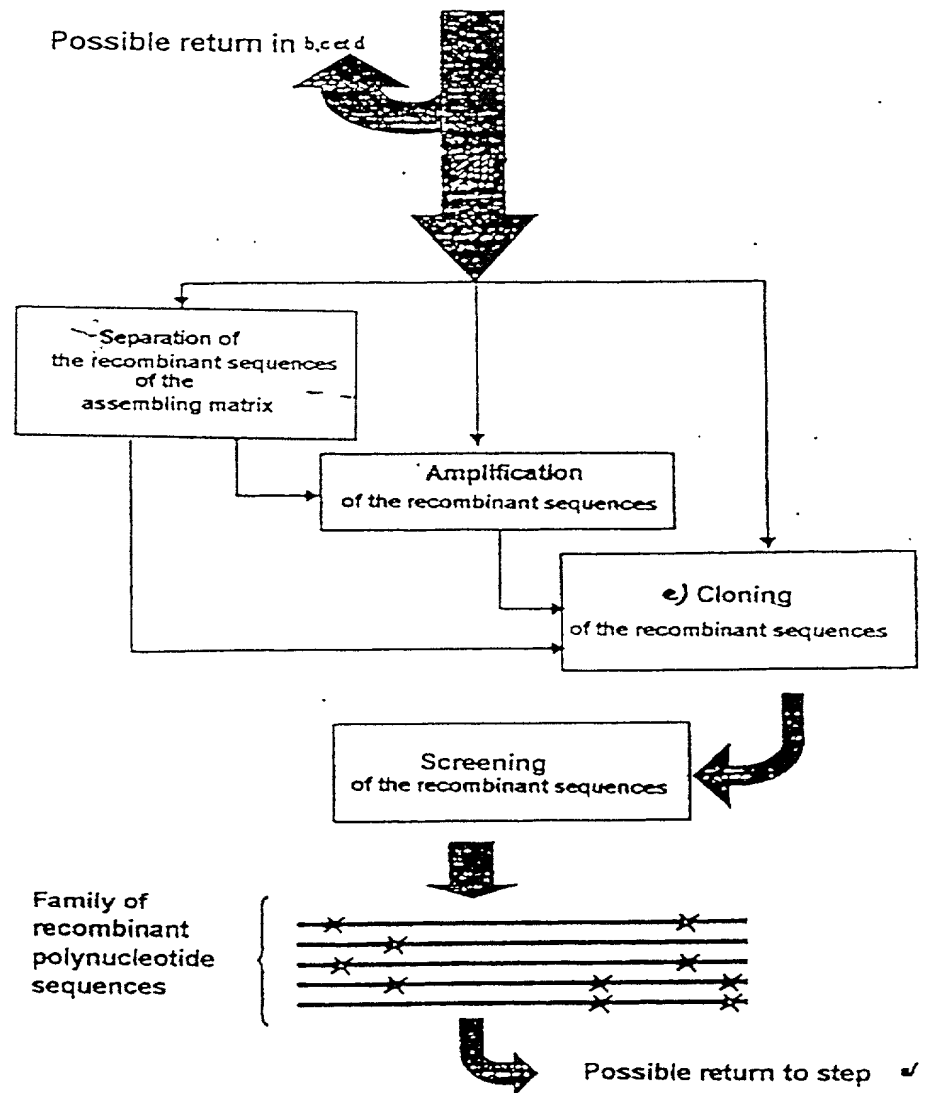


Fig.2 continued



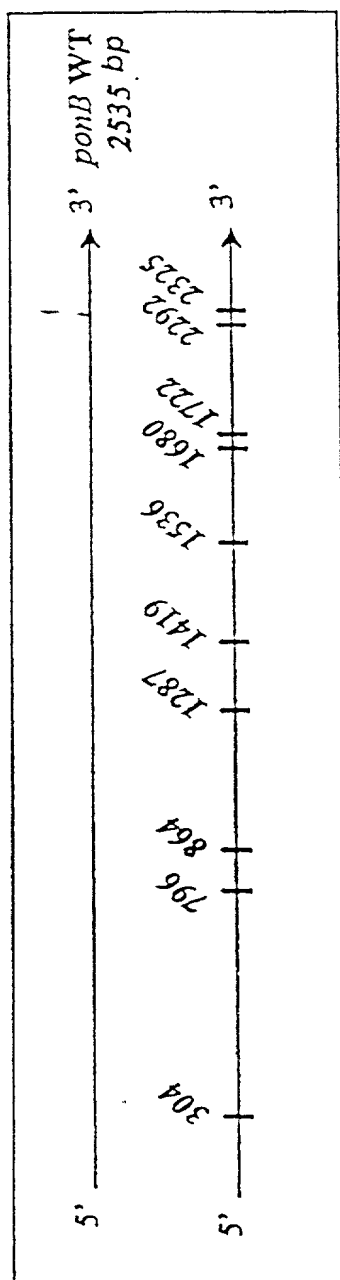


Figure 3: Position of the ten mutation zones (sites *Pvu* II and *Pst* I)

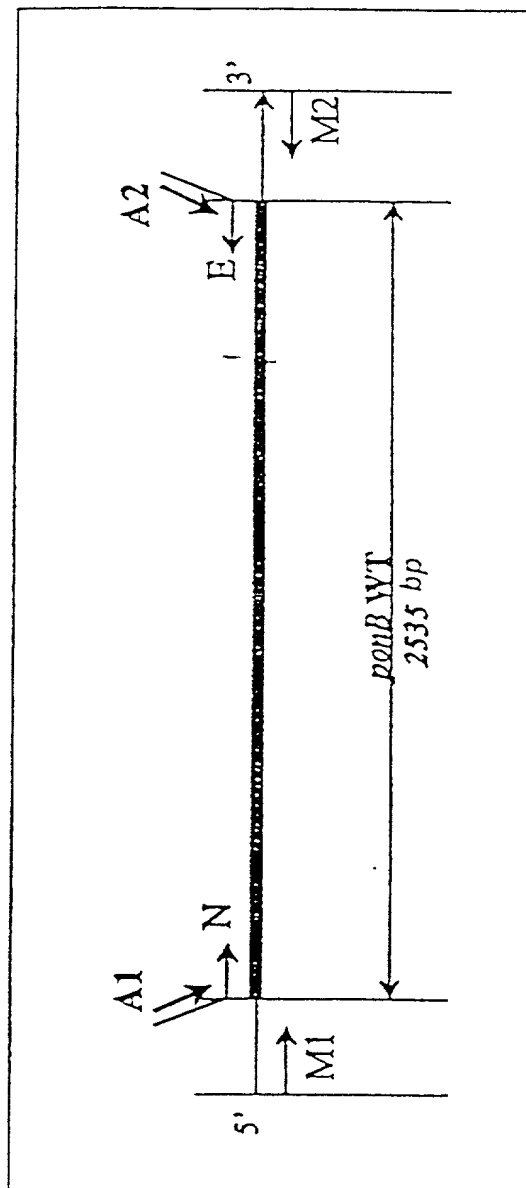


Figure 4 : Position of the primers used as compared to the sequence of the *ponB* gene

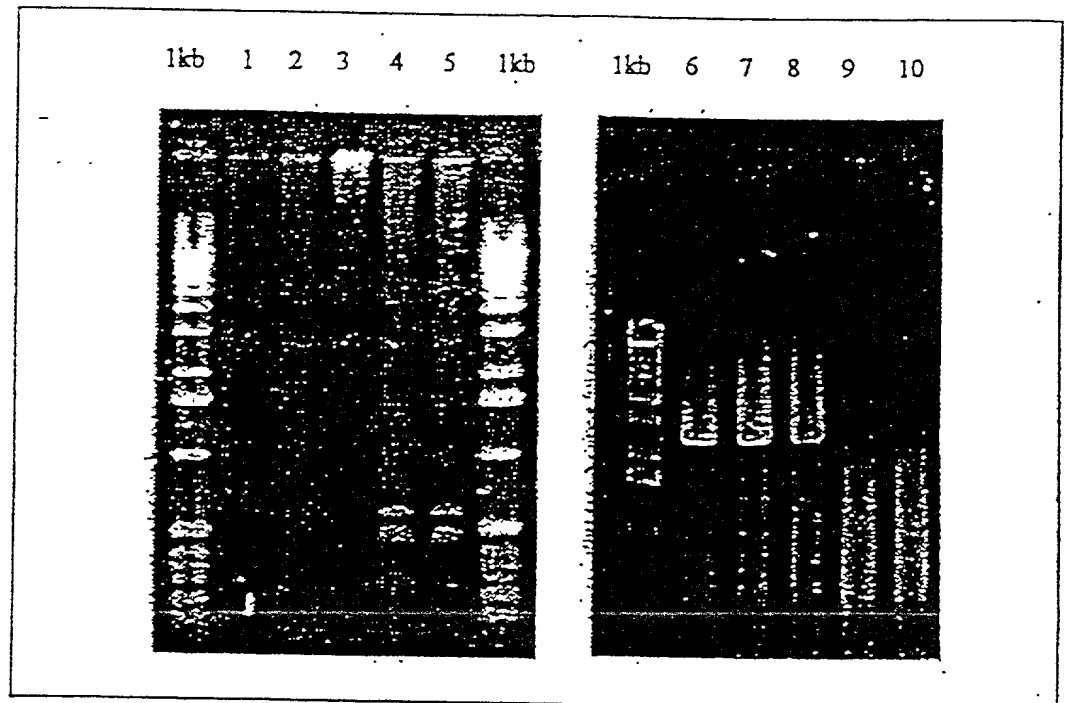


Fig. 5 : Migration of the RLR reactions and of the PCR amplifications of these reactions

Tracks:	1/RLR 1	6/PCR RLR 1
	2/RLR 2	7/PCR RLR 2
	3/RLR 3	8/PCR RLR 3
	4/RLR 4	9/PCR RLR 4
	5/RLR Control	10/PCR RLR Control

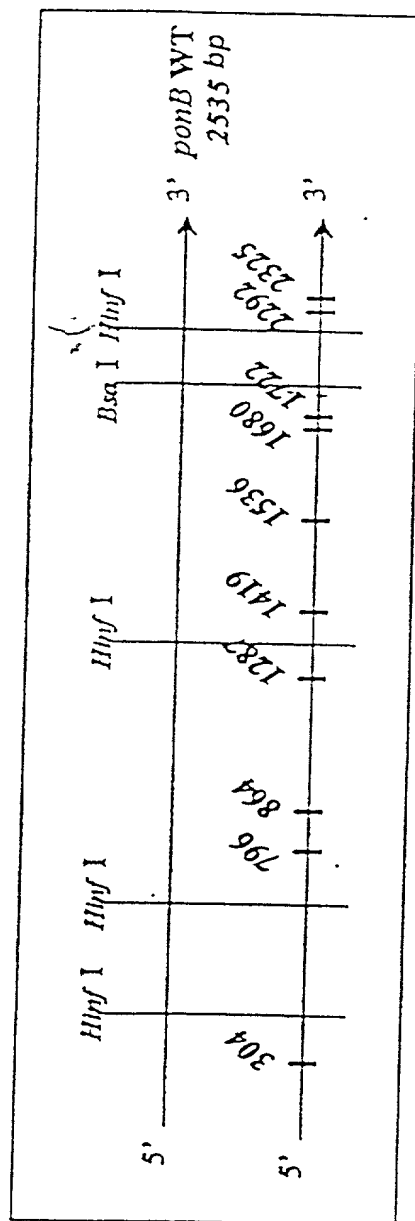


Figure 6 : Position of the mutations as compared to the restriction fragments